

PCT

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,243A

DATE: 04/15/2003 TIME: 14:01:41

Input Set : A:\09701243.txt

Output Set: N:\CRF4\04152003\I701243A.raw

3 <110> APPLICANT: MOUGIN, Bruno LAAYOUN, Ali 6 <120> TITLE OF INVENTION: METHOD FOR AMPLIFYING AT LEAST ONE SPECIFIC NUCLEOTIDE SEQUENCE, AND 7 PRIMERS USED 9 <130> FILE REFERENCE: 107976 11 <140> CURRENT APPLICATION NUMBER: US 09/701,243A 12 <141> CURRENT FILING DATE: 2000-12-28 14 <150> PRIOR APPLICATION NUMBER: PCT/FR99/01247 15 <151> PRIOR FILING DATE: 1999-05-27 17 <150> PRIOR APPLICATION NUMBER: FR 98/06866 18 <151> PRIOR FILING DATE: 1998-05-27 20 <160> NUMBER OF SEQ ID NOS: 29 22 <170> SOFTWARE: PatentIn version 3.1 24 <210> SEQ ID NO: 1 25 <211> LENGTH: 23 26 <212> TYPE: DNA 27 <213> ORGANISM: artificial sequence 29 <220> FEATURE: 30 <223> OTHER INFORMATION: Blocking primer 32 <400> SEQUENCE: 1 23 33 atccttcgtg tccccacagc acg 36 <210> SEQ ID NO: 2 37 <211> LENGTH: 19 38 <212> TYPE: DNA 39 <213> ORGANISM: artificial sequence 41 <220> FEATURE: 42 <223> OTHER INFORMATION: Blocking primer 44 <400> SEQUENCE: 2 19 45 tegeegetge actgtgaag 48 <210> SEQ ID NO: 3 49 <211> LENGTH: 24 50 <212> TYPE: DNA 51 <213> ORGANISM: artificial sequence 53 <220> FEATURE: 54 <223> OTHER INFORMATION: Blocking primer 56 <220> FEATURE: W--> 57 <221> NAME/KEY: particular characteristic 58 <222> LOCATION: (24)..(24) 59 <223> OTHER INFORMATION: modification by the C6-NH2 group 62 <400> SEQUENCE: 3 24 63 cccccaqca cgtttcttgg agct 66 <210> SEQ ID NO: 4

67 <211> LENGTH: 24

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24

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- 68 <212> TYPE: DNA
- 69 <213> ORGANISM: artificial sequence
- 71 <220> FEATURE:
- 72 <223> OTHER INFORMATION: Blocking primer
- 74 <220> FEATURE:
- W--> 75 <221> NAME/KEY: particular characteristic
 - 76 <222> LOCATION: (1)..(1)
 - 77 <223> OTHER INFORMATION: modification by the acridine group
 - 80 <220> FEATURE:
- W--> 81 <221> NAME/KEY: particular characteristic
 - 82 <222> LOCATION: (24)..(24)
 - 83 <223> OTHER INFORMATION: modification by H
 - 86 <400> SEQUENCE: 4
 - 87 cccccaqca cgtttcttgg agct
 - 90 <210> SEQ ID NO: 5
 - 91 <211> LENGTH: 26
 - 92 <212> TYPE: DNA
 - 93 <213> ORGANISM: artificial sequence
 - 95 <220> FEATURE:
 - 96 <223> OTHER INFORMATION: Blocking primer
 - 98 <220> FEATURE:
 - 99 <221> NAME/KEY: misc_feature
 - 100 <222> LOCATION: (24)..(24)
 - 101 <223> OTHER INFORMATION: n = inosine
 - 104 <220> FEATURE:
- W--> 105 <221> NAME/KEY: particular characteristic
 - 106 <222> LOCATION: (26)..(26)
 - 107 <223> OTHER INFORMATION: modification by the C6-NH2 group
 - 110 <400> SEQUENCE: 5
- 26 W--> 111 cccacagcac gtttcttgga gcangc
 - 114 <210> SEO ID NO: 6
 - 115 <211> LENGTH: 21
 - 116 <212> TYPE: DNA
 - 117 <213> ORGANISM: artificial sequence
 - 119 <220> FEATURE:
 - 120 <223> OTHER INFORMATION: Blocking primer
 - 122 <400> SEQUENCE: 6
 - 123 cccagcacgt ttcttggagc t
 - 126 <210> SEQ ID NO: 7
 - 127 <211> LENGTH: 24
 - 128 <212> TYPE: DNA
 - 129 <213> ORGANISM: artificial sequence
 - 131 <220> FEATURE:
 - 132 <223> OTHER INFORMATION: Blocking primer
 - 134 <400> SEQUENCE: 7
 - 135 cccccaqca cgtttcttgg agct
 - 138 <210> SEQ ID NO: 8
 - 139 <211> LENGTH: 24
 - 140 <212> TYPE: DNA

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141 <213> ORGANISM: artificial sequence 143 <220> FEATURE: 144 <223> OTHER INFORMATION: Blocking primer 146 <220> FEATURE: 147 <221> NAME/KEY: misc_feature 148 <222> LOCATION: (23)..(23) 149 <223> OTHER INFORMATION: n = inosine 152 <400> SEQUENCE: 8 24 W--> 153 cccccagca cqtttcttgg agnt 156 <210> SEO ID NO: 9 157 <211> LENGTH: 24 158 <212> TYPE: DNA 159 <213> ORGANISM: artificial sequence 161 <220> FEATURE: 162 <223> OTHER INFORMATION: Blocking primer 164 <220> FEATURE: 165 <221> NAME/KEY: misc_feature 166 <222> LOCATION: (22)..(22) 167 <223> OTHER INFORMATION: n = inosine170 <220> FEATURE: 171 <221> NAME/KEY: misc feature 172 <222> LOCATION: (23)..(23) 173 <223> OTHER INFORMATION: n = inosine 176 <400> SEQUENCE: 9 24 W--> 177 catttcctca atgggacgga gnna 180 <210> SEQ ID NO: 10 181 <211> LENGTH: 26 182 <212> TYPE: DNA 183 <213> ORGANISM: artificial sequence 185 <220> FEATURE: 186 <223> OTHER INFORMATION: Blocking primer 188 <220> FEATURE: 189 <221> NAME/KEY: misc_feature 190 <222> LOCATION: (24)..(24) 191 <223> OTHER INFORMATION: n = inosine 194 <400> SEQUENCE: 10 26 W--> 195 cccccagcac gtttcttgga gcangc 198 <210> SEQ ID NO: 11 199 <211> LENGTH: 26 200 <212> TYPE: DNA 201 <213> ORGANISM: artificial sequence 203 <220> FEATURE: 204 <223> OTHER INFORMATION: Blocking primer 206 <220> FEATURE: 207 <221> NAME/KEY: misc_feature 208 <222> LOCATION: (24)..(24) 209 <223> OTHER INFORMATION: n = inosine 212 <400> SEQUENCE: 11 26

W--> 213 cccacagcac gtttcttgga gcangc

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Input Set : A:\09701243.txt

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     219 <213> ORGANISM: artificial sequence
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     224 <400> SEQUENCE: 12
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     225 cacgtttctt gcagcagga
     228 <210> SEQ ID NO: 13
     229 <211> LENGTH: 22
     230 <212> TYPE: DNA
     231 <213> ORGANISM: artificial sequence
     233 <220> FEATURE:
     234 <223> OTHER INFORMATION: Blocking primer
     236 <400> SEQUENCE: 13
                                                                                 22
     237 cagcacgttt cttgcagcag ga
     240 <210> SEQ ID NO: 14
     241 <211> LENGTH: 19
     242 <212> TYPE: DNA
     243 <213> ORGANISM: artificial sequence
     245 <220> FEATURE:
     246 <223> OTHER INFORMATION: Blocking primer
     248 <220> FEATURE:
     249 <221> NAME/KEY: misc_feature
     250 <222> LOCATION: (3)..(3)
     251 <223> OTHER INFORMATION: n = inosine
     254 <400> SEQUENCE: 14
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     258 <210> SEQ ID NO: 15
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     264 <223> OTHER INFORMATION: Blocking primer
     266 <220> FEATURE:
     267 <221> NAME/KEY: misc feature
     268 <222> LOCATION: (6)..(6)
     269 <223> OTHER INFORMATION: n = inosine
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                                                                                 22
W--> 273 cagcangttt cttgcagcag ga
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     278 <212> TYPE: DNA
     279 <213> ORGANISM: artificial sequence
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     284 <220> FEATURE:
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     286 <222> LOCATION: (10)..(10)
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Input Set : A:\09701243.txt

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	287	<223>	OTHER INFORMATION: n = inosine	
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			SEQ ID NO: 17	
			LENGTH: 26	
			TYPE: DNA	
			ORGANISM: artificial sequence	
			FEATURE:	
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	303	<221>	NAME/KEY: misc_feature	
	304	<222>	LOCATION: (10)(10)	
	305	<223>	OTHER INFORMATION: n = inosine	
			SEQUENCE: 17	
M>	309 cccacagcan gtttcttgca gcagga			26
			SEQ ID NO: 18	
			LENGTH: 26	
			TYPE: DNA	
			ORGANISM: artificial sequence	
			FEATURE:	
			OTHER INFORMATION: Blocking primer	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: (10)(10) OTHER INFORMATION: n = inosine	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: (25)(25)	
			OTHER INFORMATION: n = inosine	
			SEQUENCE: 18	
W>			agcan gtttcttgca gcagna	26
			SEQ ID NO: 19	
			LENGTH: 26	
	338	<212>	TYPE: DNA	
	339	<213>	ORGANISM: artificial sequence	
			FEATURE:	
			OTHER INFORMATION: Blocking primer	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: (10)(10)	
			OTHER INFORMATION: n = inosine	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: (25)(25)	
			OTHER INFORMATION: n = inosine	
T.T >			SEQUENCE: 19	26
W>			agcan gtttettgea geagna SEQ ID NO: 20	20
			LENGTH: 33	
	201	\ 211>	DENGIN. JJ	

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/701,243A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 24 Seq#:8; N Pos. 23 Seq#:9; N Pos. 22,23 Seq#:10; N Pos. 24 Seq#:11; N Pos. 24 Seq#:14; N Pos. 3 Seq#:15; N Pos. 6 Seq#:16; N Pos. 10 Seq#:17; N Pos. 10 Seq#:18; N Pos. 10,25 Seq#:19; N Pos. 10,25

VERIFICATION SUMMARY

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Input Set : A:\09701243.txt

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L:57 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:75 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:81 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:105 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:195 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:291 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:333 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0